

SEQUENCE LISTING

DEN 110> DARNAY, BRYANT G.

<120> METHODS AND COMPOSITIONS USING POLYNUCLEOTIDES AND POLYPEPTIDES OF RANK-ASSOCIATED INHIBITOR (RAIN)

<130> UTSC:761US

<140> 10/629,329

<141> 2003-07-29

<150> 60/399,205

<151> 2002-07-29

<160> 23

<170> PatentIn Ver. 2.1

<210> 1

<211> 729

<212> DNA

<213> Homo sapiens

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<221> CDS

<222> (1)..(729)

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1 10 15

ggc gcg cag gac aag gag cat cca aga tac ctg atc cca gaa ctt tgc 96
Gly Ala Gln Asp Lys Glu His Pro Arg Tyr Leu Ile Pro Glu Leu Cys
20 25 30

aaa cag ttt tac cat tta ggc tgg gtc act ggg act gga gga gga att 144 Lys Gln Phe Tyr His Leu Gly Trp Val Thr Gly Thr Gly Gly Gly Ile 35 40 45

agc ttg aag cat ggc gat gaa atc tac att gct cct tca gga gtg caa 192 Ser Leu Lys His Gly Asp Glu Ile Tyr Ile Ala Pro Ser Gly Val Gln 50 55 60

aag gaa cga att cag cct gaa gac atg ttt gtt tgt gat ata aat gaa 240 Lys Glu Arg Ile Gln Pro Glu Asp Met Phe Val Cys Asp Ile Asn Glu 65 70 75 80

_	_		_			tcg Ser		_	_	_				_	_	288
_					_	aat Asn	_			_	_		_		-	336
						aaa Lys	_	_		_	_					384
						att Ile 135					_					432
						999 999										480
_						aca Thr								_	_	528
						gaa Glu			-		_	_	_	_	_	576
	-			_		gtg Val			_				_	-		624
	_	_		_		gac Asp 215				_		-	_		_	672
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Gly Ala Gln Asp Lys Glu His Pro Arg Tyr Leu Ile Pro Glu Leu Cys
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Lys Gln Phe Tyr His Leu Gly Trp Val Thr Gly Thr Gly Gly Gly Ile
                             40
Ser Leu Lys His Gly Asp Glu Ile Tyr Ile Ala Pro Ser Gly Val Gln
                         55
Lys Glu Arg Ile Gln Pro Glu Asp Met Phe Val Cys Asp Ile Asn Glu
                     70
Lys Asp Ile Ser Gly Pro Ser Pro Ser Lys Lys Leu Lys Lys Ser Gln
                                     90
Cys Thr Pro Leu Phe Met Asn Ala Tyr Thr Met Arg Gly Ala Gly Ala
                                105
Val Ile His Thr His Ser Lys Ala Ala Val Met Ala Thr Leu Leu Phe
                            120
Pro Gly Arg Glu Phe Lys Ile Thr His Gln Glu Met Ile Lys Gly Ile
                        135
Lys Lys Cys Thr Ser Gly Gly Tyr Tyr Arg Tyr Asp Asp Met Leu Val
                    150
                                        155
Val Pro Ile Ile Glu Asn Thr Pro Glu Glu Lys Gly Leu Lys Asp Arg
                                    170
Met Ala His Ala Met Asn Glu Tyr Pro Asp Ser Cys Ala Val Leu Val
                                185
Arg Arg His Gly Val Tyr Val Trp Gly Glu Thr Trp Glu Lys Ala Lys
                            200
Thr Met Cys Glu Cys Tyr Asp Tyr Leu Phe Asp Ile Ala Val Ser Met
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                                            220
Lys Lys Val Gly Leu Asp Pro Ser Gln Leu Pro Val Gly Glu Asn Gly
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Ile Val

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								cga Arg		_			_		_		96	
								gtc Val 40								_	144	
								tac Tyr								_	192	
								atg Met			-	_				_	240	
·	-		_				_	tct Ser	_	_	_			_	_	_	288	
								tat Tyr									336	
								gct Ala 120						_			384	
								cat His			_						432	
								tac Tyr	_		_	_	_			_	480	
								gaa Glu									528	
								cca Pro			_		_		_		576	
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Met Cys Glu Cys Tyr Asp Tyr Leu Phe Asp Ile Ala Val Ser Met Lys
    210
                        215
aag atg gga ctc gat cca aca caq ctc cca qtt qqa qaa aat qqa att
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Lys Met Gly Leu Asp Pro Thr Gln Leu Pro Val Gly Glu Asn Gly Ile
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Leu Lys His Gly Asn Glu Ile Tyr Ile Ala Pro Ser Gly Val Gln Lys
                         55
                                             60
Glu Arg Ile Gln Pro Glu Asp Met Phe Val Cys Asp Ile Asn Glu Gln
                     70
                                         75
Asp Ile Ser Gly Pro Pro Ala Ser Lys Lys Leu Lys Lys Ser Gln Cys
                 85
                                     90
Thr Pro Leu Phe Met Asn Ala Tyr Thr Met Arg Gly Ala Gly Ala Val
                                105
Ile His Thr His Ser Lys Ala Ala Val Met Ala Thr Leu Leu Phe Pro
                            120
Gly Gln Glu Phe Lys Ile Thr His Gln Glu Met Ile Lys Gly Ile Arg
                        135
Lys Cys Thr Ser Gly Gly Tyr Tyr Arg Tyr Asp Asp Met Leu Val Val
                    150
                                        155
Pro Ile Ile Glu Asn Thr Pro Glu Glu Lys Asp Leu Lys Glu Arg Met
                165
                                    170
Ala His Ala Met Asn Glu Tyr Pro Asp Ser Cys Ala Val Leu Val Arg
                                185
Arg His Gly Val Tyr Val Trp Gly Glu Thr Trp Glu Lys Ala Lys Thr
                            200
Met Cys Glu Cys Tyr Asp Tyr Leu Phe Asp Ile Ala Val Ser Met Lys
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210 215 220

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Ser Cys

<210> 6

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Lys Thr Leu Ala Asp Ala Met Ala His Thr Met Asn Glu Tyr Pro Asp 20 25 30

Ser Cys

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C1007 22
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- · · · · · · · · · · · · · · · · · · ·
Primer
Primer

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